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Minimum
Maximum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           n DB seq
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Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2000 Compugen Ltd.
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Ewe colostrinin pe
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S. pneumoniae elon
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Colostrinin peptid
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-	Human polypeptide Rat GABA-A recepto Novel human diagno	onibacterium	SB5	2 strain SB5	2 strain SB5	HSV-2 strain sas C	2 strain SB5	2 strain SB5	2 strain SB5	strain sas	# 8540deu		E-prototo	lipovin ad		G Proceru	orbuan e i	nepckse #	nGPCR58 #	beta-s	<u>1e1</u>	otein.	#1970	de l	de #1973	bone marrow	rain	1n #1954	Peptide #2052 enco

ALIGNMENTS

RESULT 1 AAB72247 PXXXXX 17-AUG-1999; 17-AUG-2000; 2000WO-US22818 WO200111937-A2. Colostrinin; immune response; cytokine; blood cell proliferation; central nervous system disorder; neurological diosrder; mental di 14-MAY-2001 (first entry) AAB72247 standard; peptide; 17 (TEXA) UNIV TEXAS SYSTEM. (REGE-) REGEN THERAPEUTICS PLC. 22-FEB-2001 Synthetic. neurosis; Colostrinin derived cytokine inducing peptide SEQ ID AAB72247; infection neurodegenerative disease; Alzheimer's disease; 9908-0149311. æ

Inducing a cytokine and modulating an immune response, useful for treating central nervous system diseases and bacterial and viral

WPI; 2001-202804/20.

Stanton GJ,

Hughes TK,

Boldogh I,

Georgiades

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RESULT
AAB72501
ID AAB;
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AC AAB;
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Best Local Similarity 100
Matches 17; Conservative
                         The present invention relates to a method for modulating the oxidative stress level in a cell or a patient, comprising contacting the cell with, or administering to the patient, an oxidative stress regulator selected from colostrinin, or its constituent peptide (e.g. the present peptide), to change the level of an oxidating species in the cell. The method can be used to treat oxidative damage to skin, by decreasing or preventing an increase in the level of damage to a biomolecule of the patient.
                                                                                                                                          Claim 6;
                                                                                                                                                                 Modulating oxidative stress level in a cell, involves cell with an oxidative stress regulator selected from constituent peptide, analog or their combinations
                                                                                                                                                                                                                              PI;
                                                                                                                                                                                                                                                         Stanton
                                                                                                                                                                                                                                                                                  (TEXA ) UNIV
                                                                                                                                                                                                                                                                                                              17-AUG-1999;
                                                                                                                                                                                                                                                                                                                                           17-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Colostrinin peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                        22-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                   WO200112650-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB72501;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB72501 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             inducing cytokines. Colostrinin and its derived peptides are useful for inducing cytokine production, for modulating an immunological response and for inducing blood cell proliferation. The peptides are useful in treatment of disorders of the central nervous system, neurological disorders, mental disorders, dementia, neurodegenerative diseases, also included the contral disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          acquired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Alzheimer's disease, motor neurone disease, disorders of the immune system, bacterial ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      peptides have immune response modulatory activity, and are capable of inducing cytokines. Colostrinin and its derived peptides are useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequences AAB72246 - AAB72275 represent peptides derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infections, regulator -
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                                                                                                                                                                                                                           2001-218342/22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOTPOPLLOVMMEPOGD 17
                                                                                                                                                                                                                                                       GJ,
                                                                                                                                          Page 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of the immune system, bacterial and viral infections and
immunological deficiencies.
17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       comprises administering colostrinin as
                                                                                                                                                                                                                                                                                                                                           2000WO-US22665
                                                                                                                                                                                                                                                                                    TEXAS SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                       Hughes
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                                                                                                                                                                                                                                                                                                              9905-0149310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polypeptide aggregate contained in colostrum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         oxidative stress regulator;
                                                                                                                                        48pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             50pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Peptide;
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                                                                                                                                                                                                                                                       Boldogh I;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         colostrinin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   psychosis,
                      decreasing or preventing ule of the patient.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       an immunological
                                                                                                                                                                                contacting the colostrinin, its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 clostrinin,
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Sequence

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Sheep; colostrinin; proline rich polypeptide; colostrum;

immune disorder;

fragment A-1.

Ewe colostrinin 21-MAR-2001

(first

entry)

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AAB59310
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Matches 17
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Best Local
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                                                                                                                                                                                        Sequence
                                                                                                                                                                                                        The present invention relates to a method for promoting neural cell differentiation and treating damaged neural cells, using colostrinin and colostrinin constituent peptides (e.g. the present peptide) as a neural cell regulator. Colostrinin is a polypeptide complex found in colostrum
                                                                                                                                                                                                                                                                     Use of colostrinin, its constituent peptide or analog as a neural regulator, for promoting neural cell differentiation and treating damaged neural cells in a patient -
                                                   AAB59310;
                                                                   AAB59310 standard;
                                                                                                                                                                                                                                                    Claim 6; Page 21; 35pp; English
                                                                                                                                                                                                                                                                                                                        Boldogh
                                                                                                                                                                                                                                                                                                                                                           17-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                colostrum.
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                                                                                                                                                       Similarity
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                                                                                                                                                Conservative
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                                                                   Peptide;
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1.2e-07;
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disease, Alzhelmer's disease, psychosis and neurosis, immune system disorders such as bacterial and viral infections, to improve the development of a child's immune system, as a dietary supplement, and promote the dissolution of beta-amyloid plaques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides the sequences of a number of peptides found in ewe's colostrinin. Colostrinin is the proline-rich polypeptide fragment of colostrum. These peptides can be used in the treatment of central nervous system disorders such as senile dementia, Parkinson's
                                                                                                                                                      Ovis sp
                                                                                                                                                                                                             Ewe colostrinin peptide fragment derived sequence #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ving an N-terminal amino acid sequence isolated from and immune system, viral and bacterial infections, and diseases characterized by amyloid plaques
                                (REGE-) REGEN THERAPEUTICS PLC
                                                        02-JUN-1999;
                                                                               02-JUN-2000; 2000WO-GB02128
                                                                                                         14-DEC-2000
                                                                                                                               WO200075173-A2
                                                                                                                                                                              central
                                                                                                                                                                                          Sheep; colostrinin;
                                                                                                                                                                                                                                          21-MAR-2001
                                                                                                                                                                                                                                                                  AAB59341;
                                                                                                                                                                                                                                                                                        AAB59341 standard; Peptide; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 7; Page 27; 63pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-JUN-2000; 2000WO-GB02128
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (REGE-) REGEN THERAPEUTICS PLC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  central nervous
                                                                                                                                                                                                                                                                                                                                                                                                   Match 100.0%;
Local Similarity 100.0%;
les 17; Conservative (
                                                                                                                                                                                                                                                                                                                                                     ü
                                                                                                                                                                                                                                                                                                                                                                           1 LOTPOPLLOVMMEPOGD 17
                                                                                                                                                                                                                                                                                                                  G
                                                                                                                                                                                                                                                                                                                                                   ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ~~1-071058/08
                                                                                                                                                                              nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                               17
                                                                                                                                                                                                                                          (first entry)
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                                                        99GB-0012852
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                                                                                                                                                                             system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             system disorder; dietary supplement;
                                                                                                                                                                            proline rich
tem disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                              Score 89;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                         polypeptide; colostrum; immune disorder;
dietary supplement; beta-amyloid plaque.
                                                                                                                                                                                                                                                                                                                                                                                                             1.2e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                        22;
                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                        Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               tral nervous system and diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             beta-amyloid plaque.
                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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The invention relates to colostrinin peptide fragments which are useful, inter alia, in the treatment of chronic disorders of the immune system and the central nervous system. Colostrinin peptides are used as a

Peptide useful as an inter alla in the treatment of e.g. disorders of the immune system and the central nervous system comprises ten amino-terminal amino acid sequence derived from peptides present in colostrinin -

WPI; 2001-488775/53.

26-JAN-2000; 2000GB-0001825. 26-JAN-2001; 2001WO-GB00329

(REGE-) REGEN

THERAPEUTICS PLC

02-AUG-2001 WO200155199-A1. Unidentified.

Claim 1;

Page 15; 40pp; English

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RESULT
AAE07185
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                        Colostrinin; nootropic; neuroprotective; immunomodulatory; antibacterial; Parkinson's disease; Alzheimer's disease; mental disorder; food additive; central nervous system disorder; neurodesgenerative disorder; weight loss; beta amyloid plaque; psychosis; neurosis; cachexia; motor neuron disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides the sequences of a number of peptides found in ewe's colostrinin. Colostrinin is the proline-rich polypeptide fragment of colostrum. These peptides can be used in the treatment of central nervous system disorders such as senile dementia, Parkinson's disease, Alzheimer's disease, psychosis and neurosis, immune system disorders such as bacterial and viral infections, to improve the development of a child's immune system, as a dietary supplement, and to
                                                                                                                                                                                                                                                                                                                                                                acquired
                                                                                                                                                                                                                                                                                                                                                                                                                                            Colostrinin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAE07185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAE07185 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                promote
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 8; Page 27; 63pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Peptides having an N-terminal amino acid sequence isolated from colostrinin for treating e.g. disorders of the central nervous and immune system, viral and bacterial infections, and diseases characterized by amyloid plaques -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-071058/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 LOTPOPLLOVMMEPOGD 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 lqtpqpllqvmmepqgd 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the dissolution of beta-amyloid plaques
                                                                                                                                                                                                                                                                                                                                                               immunological deficiency;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 89;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                               neurological disorder; dementia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                system
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133,3424,24
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AAE07195
ID AAE0
XX AAE0
AC AAE0
XX O6-N
XX Mod1
VX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Color des
                                                      Peptide useful as an inter alia in the treatment of e.g. disorders of the immune system and the central nervous system comprises ten amino-terminal amino acid sequence derived from peptides present in colostrinin.
                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                          Georgiades
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             adults and senile persons, who have been subjected to chemotherapy or have suffered from cachexia or weight loss due to the chronic disease. Colostrinin peptides are also used as food additives and as an auxillary withdrawal treatment for drug addicts, after a period of detoxification and in persons dependent on stimulants. Colostrinin peptides are used to prepare antibodies and to treat emotional disturbances, e.g. emotional disturbances of psychiatric patients in a state of depression. These
                                                                                                                                                                                                                                                                                                                                                                          26-JAN-2000;
                                                                                                                                                                                                                                                                                                              (REGE-) REGEN THERAPEUTICS PLC
                                                                                                                                                                                                                                                                                                                                                                                                                             26-JAN-2001; 2001WO-GB00329.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Modified colostrinin cyclic peptide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAE07195 standard; peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 colostrinin peptides improves the development of immune system in a new born child and to correct the immunological deficiencies in a child. The present sequence is colostrinin peptide 1 related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          medicament in the treatment of neurological disorders e.g., dementia, neurodegenerative disorders e.g., Alzheimer's disease, motor neuron disease e.g., Parkinson's disease, mental disorders e.g. psychosis and neurosis, in acquired immunological deficiencies, chronic bacterial and viral infections and diseases characterised by the presence of beta-amyloid plaques and as a dietary supplement for babies, small children,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAE07195
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mor. 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100 nes 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 QPLLQVMMEPQ 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 qpllqvmmepq 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     strinin; nootropic; neuroprotective; immunomodulatory; inson's disease; Alzheimer's disease; mental disorder;
                                                                                                                                                                                              2001-488775/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'sl nervous system disorder; sdisease; mental disorder; food additive; myloid plaque; psychosis; neurodegenerative disorder; weight loss; cyd immunological deficiency; neurological disorder; disor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   J199-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note= "N-terminal acetyl; this residue forms
.inkage with Gln found at the C-terminal end"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 62.9%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ξ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 22;
0.014;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Example 2; Page

8; 40pp; English

Claim 3; Pages 312-314; 333pp; English

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AAW80722
ID AAW8
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Best Local :
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                                                                                                                                          Baltz RH, Bur
Mills BJ, Nor
Skatrud PL, Si
Young Bellido
                                        Streptococcus pneumoniae nucleic acid sequences - used in DNA chips for evaluating gene expression, and identification of virulence
                                                                                            WPI; 1998-348529/30.
N-PSDB; AAV65279.
                                                                                                                                                                                                                                                                                                                                                                                                                               Streptococcus pneumoniae protein; recombinant; gene expression; DNA chip; virulence; antibody; infection; detection; treatment; hypothetical;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   viral infections and diseases characterised by the presence of beta amyloid plaques and as a dietary supplement for babies, small children, adults and senile persons, who have been subjected to chemotherapy or have suffered from cachaxia or weight loss due to the chronic disease. Colostrinin peptides are also used as food additives and as an auxillary withdrawal treatment for drug addicts, after a period of detoxification and in persons dependent on stimulants. Colostrinin peptides are used to prepare antibodies and to treat emotional disturbances, e.g. emotional disturbances of psychiatric patients in a state of depression. These
                                                                                                                                                                                                                                                                                         09-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                    cell wall biosynthetic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       S. pneumoniae
                                                                                                                                                                                                                                                          13-DEC-1996;
                                                                                                                                                                                                                                                                                                                         18-JUN-1998
                                                                                                                                                                                                                                                                                                                                                         WO9826072-A1
                                                                                                                                                                                                                                                                                                                                                                                     Streptococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAW80722;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW80722 standard;
                                                                                                                                                                                                                          (ELIL ) LILLY & CO ELI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             colostrinin peptides improves the development of immune system in a new
born child and to correct the immunological deficiencies in a child.
The present sequence is modified colostrinin cyclic peptide #1 related to
the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to colostrinin peptide fragments which are useful, inter alia, in the treatment of chronic disorders of the immune system and the central nervous system. Colostrinin peptides are used as a medicament in the treatment of neurological disorders e.g., dementia, neurodegenerative disorders e.g., Alzheimer's disease, motor neuron
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               neurosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 QPLLQVMMEPQ 15
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                                                                                                                                          Burgett SG,
Norris FH,
L, Smith MC,
lido ML;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     g., Parkinson's disease, mental disorders e.g. psychosis and in acquired immunological deficiencies, chronic bacterial and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62.9%;
ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                     pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    elongation factor G.
                                                                                                                                                                                                                                                         96US-0036281
                                                                                                                                                                                                                                                                                         97WO-US22578.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein; 693
                                                                                                                                                       Dehoff BS, Hoskins JA, Jaskunas SR;
Peery RB, Rockey PK, Rosteck PR;
Solenberg PJ, Treadway PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              p; Score 56; DB
p; Pred. No. 0.0
p; Mismatches
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0.016;
                                                                                                                                                                                                                                                                                                                                                                                                                  minimal gene set
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                                                                                                                                                                                                                                                                                                                                                                                                                  protein.
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AAU37640
ID AAU3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequences (AAW80605 to AAW80728). The protein sequences are classified as hypothetical, cell wall biosynthetic, external target, or minimal generate proteins. A recombinant host containing a vector comprising any of the above nucleic acids can be used for the recombinant expression of the proteins. The invention also provides a DNA chip having arrayed on it at the DNA chip can be used methods for evaluating gene expression in some phenomena and for identifying virulence genes in S. pneumoniae and for identifying virulence genes in S. pneumoniae that selectively bind to the above proteins or peptide fragments can be used to treat S. pneumoniae infection. The antibodies that selectively bind to the above proteins or peptide
                                                                     New polynucleotides for antibiotics, comprise se
                                                                                                            N-PSDB; AAS55499.
                                                                                                                      WPI; 2001-611495/70
                                                                                                                                               Haselbeck R,
Yamamoto RT,
                                                                                                                                                                                                                                                                                                           21-MAR-2001; 2001WO-US09180.
                                                                                                                                                                                                                                                                                                                                                              WO200170955-A2
                                                                                                                                                                                                                                                                                                                                                                                     Streptococcus pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                       Antisense; prokaryotic cellular proliferation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU37640;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAU37640 standard; Protein; 693 AA
                                                                                                                                                                                                                                   27-NOV-2000;
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                                                                                                                                                                                  (ELIT-) ELITRA PHARM
                                                                                                                                                                                                             16-FEB-2001;
                                                                                                                                                                                                                                                   23-OCT-2000;
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les 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                               antibacterial;
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                                                                                                                                                                                                                                               2000US-207727P.
2000US-242578P.
                                                                                                                                                                                                                                                                                     2000US-191078P.
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                                                                                                                                             Ohlsen
Xu HH;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 pneumoniae cellular proliferation protein
                                                                                                                                                                                   INC
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                                                                 the identification and development equences of antisense nucleic acids
                                                                                                                                                         Zyskind
                                                                                                                                                                                                                                                                                                                                                                                                              drug design.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 46;
Pred. No.
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46;
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                                                                                                                                                       Trawick JD,
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The invention relates to antisense inhibitors of genes essential to prokaryotic cellular proliferation, their use in identifying the genes, their use in the discovery of novel antibiotics, the essential to

essential

The present invention relates to nucleic acids (AAH90701-AAH90918) encoding polypeptides (AAM01002-AAM01114), which are essential for the viability of a bacterial cell wall. The accompm CFE stands for "CEG For Expression", where CEG stands for "Conserved Essential Gene". The nucleic acids are useful for detecting the presence of proteins essential for the viability of a bacterial cell wall in samples such as cells, tissues,

Example 3;

Seq ID No 13233; 511pp; English

sequences

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Claim 27; Pages 344-346; 380pp; English

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RESULT
AAM0109
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Best Local
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                                                                                                                   Nucleic acids encoding conserved essential genes involved in bacterial replication which are potential targets for the treatment of antibiotic resistant bacterial infections -
                                                                                                                                                                                                                                                                                                                                                                                                                          Antibacterial; vaccine; gene therapy; bacterial cell wall viability; CFE; CEG; Conserved Essential Gene; bacterial infection; antibiotic resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 genes themselves and the encoded proteins. The prokaryotes used are Bscherichia coli, Staphylococcus aureus, Salmonella typhi, Klebsiella pneumoniae, Pseudomonas aeruginosa and Enterococcus faecalis. The invention is also useful for the identification of potential new targets for antibiotic development. The antisense nucleic acids can also be used to identify proteins used in proliferation, to express these proteins, and to obtain antibodies capable of binding to the expressed proteins. The proteins can be used to expense compounds in rational drug discovery
                                                                                                                                                                           N-PSDB;
                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                Dougherty TJ,
                                                                                                                                                                                                                                                                                                                                                                                                   Streptococcus pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           a wide variety of organisms. The present sequence represents essential prokaryotic cellular proliferation protein. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at
                                                                                                                                                                                                                                                                                        30-DEC-1999;
                                                                                                                                                                                                                                                                                                                 29-DEC-2000; 2000WO-US35604
                                                                                                                                                                                                                                                                                                                                                                         WO200149721-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CFE 96 protein sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAM01093;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAM01093 standard; Protein;
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                                                                                                                                                                                                                                                            (BRIM ) BRISTOL-MYERS SQUIBB CO
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                                                                                                                                                                           2001-496721/54.
DB; AAH90792.
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                                                                                                                                                                                                                     JA;
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                                                                                                                                                                                                                                Pucci MJ,
                                                                                                                                                                                                                                                                                       9905-0174089
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                                                                                                                                                                                                                               Dougherty BA,
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Pred. No.
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ed. No. 46;
Mismatches
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Best Local Similarity
Matches 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   and for detecting corresponding target nucleic acid molecules with complementary sequences. The nucleic acids are also useful for determining whether a genomic nucleotide sequence of interest is essential for viability of a bacterial cell or whether it resides within an operon, by integrating an exogeneous nucleotide sequence comprising a portion of an open reading frame of the genomic sequence of interest (comprising 200°500 base pairs) into the genomic sequence of interest which confers a selectable phenotype to the cell, and determining cell viability with a selectable phenotype to the cell, and determining cell cacids and proteins are also useful as vaccines and for treating bacterial infections with gene therapy and antisense therapy. The nucleic acids moleculation of targets suitable for the treatment of antibiotic resistant bacterial infections.
                                                                                                                                                                                                                                                                          Peptide #2023 encoded by breast cell single exon
                                                                                                                                                                                                                                                                                                             01-FEB-2002
                                                                                                                                                                                                                                                                                                                                         ABB29372;
                                                                                                                                                                                                                                                                                                                                                                   ABB29372 standard;
                                                                                                                                                                                                                                      disease; cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  biological fluids, and for detecting
                                                                                                                                                                                                                                                  Human; microarray;
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1 LOTPOPLLOVMMEPO
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                                       2000US-0608408
2000US-0234687.
2000US-0236359.
2000GB-0024263.
                                                                       2000US-0180312
2000US-0207456
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                                                                                                                                                                                                                                                                                                         (first
                                                                                                                   2001WO-US00662
                                                                                                                                                                                                                                                single exon probe;
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                                                                                                                                                                                                                                                                                                          entry)
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Pred.
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                                                                                                                                                                                                                                               gene expression; breast;
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46;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 693;
                                                                                                                                                                                                                                                                            nucleic acid
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CCCCXXX PTTXXXXI

WPI;

2001-496933/54.

Penn

SG,

Hanzel

R,

Chen

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Rank

(MOLE-) MOLECULAR DYNAMICS INC

Claim

27;

SEQ

IJ

NO 12340;

327pp + sequence listing;

English.

probes, human

The

New spatially-addressable set of single exon nucleic acid properties for measuring gene expression in sample derived from breast, comprises number of single exon nucleic acid probes

the

from mRNA

of human breast,

and then measuring

form

part

invention relates to a spatially-addressable set of single exon leic acid probes for measuring gene expression in a sample derived human breast and BT 474 cells. The method involves contacting probes with a collection of detectably labelled nucleic acids

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RESULT 1
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Best Local Sim
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26-MAY-2000;
30-JUN-2000;
measuring human gene expression in a sample derived from human foetal liver. The single exon nucleic acid probes may be used for predicting, measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a peptide encoded by a single exor nucleic acid probe of the invention.

Note: The sequence data for this patent did not form part of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     bound to each probe of the microarray. The probes are useful for verifying the expression of regions of genomic DNA predicted to encode proteins. They are useful for gene discovery, and for determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical agents on cells. The microarray of this invention presents a far great diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence is a peptide encoded by a single exon nucleic acid
                                                                                 The invention relates to a single exon nucleic acid
                                                                                                                Claim
                                                                                                                                                      Human
                                                                                                                                                                                 WPI;
                                                                                                                                        analyzing
                                                                                                                                                                                                            Penn
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Note: The sequence data for this patent did not form part of printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200157277-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; foetal liver; gene expression; single exon nucleic acid probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Peptide #2052
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABB34546;
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                                                                                                                                       genome-derived single exon nucleic acid probes ing gene expression in human fetal liver -
                                                                                                              27;
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9; Conserv
                                                                                                              SEQ
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                                                                                                                                                                                                                                                              2000US-0207456
2000US-0608408
2000US-0632366
2000US-0234687
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2000US-0236359
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                                                                                                          639pp + sequence listing;
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Pred. No.
                                                                                                                                                                                                            Rank
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                                                                                 probe for
                                                                                                            English.
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Best Local
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27-SEP-2000;
04-OCT-2000;
                               measuring human gene expression in a sample derived from human heart (see ABA21535-ABA41305). The present sequence is a protein encoded by one sucl probe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular system e.g. cardiovascular disease, hypertension, cardiac arrhythmias and congenital heart disease. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                    WPI;
                                                                                                                                   measuring human gene
ABA21535-ABA41305). T
                                                                                                                                                        The present invention relates to single exon nucleic acid probes for
                                                                                                                                                                                      Claim 15;
                                                                                                                                                                                                                          Single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                           04-FEB-2000;
26-MAY-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; gene expression; heart; microarray; vascular system; cardiovascular disease; hypertension; cardiac arrhythmia;
                                                                                                                                                                                                                 hearts
                                                                                                                                                                                                                                                                            Penn SG,
                                                                                                                                                                                                                                                                                                     (MOLE-)
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03-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   congenital heart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein #1954 encoded by probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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hes 9; Conservat
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2000US-0207456.
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                                                                                                                                                                                    ID No 21725; 530pp; English.
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26-MAY-2000;
30-JUN-2000;
                                                                                                                                                               The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is a protein encoded by one of
                                                                                                                                                                                                                                                                                                                                                                  Penn
                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                       Single exon nucleic acid probes for analyzing gene expression in human
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04-FEB-2000; 2000US-0180312.
26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-032366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                  The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow sar ', which may enable the improved diagnosis and treatment of cancers lymphoma, leukaemia and myeloma. The present sequence is a concern of encoded by one of the probes of the invention.
                                                                                                                                                                                                                                                                                           Example 4; SEQ ID NO: 28035; 658pp + Sequence Listing; English
                                                                                                                                                                                                                                                                                                                             Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human bone marrow
                                                                                                                                                                                                                                                                                                                                                                                                             Penn SG,
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                              GENERAL INFORMATION:
APPLICANT: Katz, Leonard
APPLICANT: Stassi, Diane L.
APPLICANT: Summers Jr., Richard G
APPLICANT: Ruan, Xiaoan
APPLICANT: Ruan, Xiaoan
APPLICANT: Kakavas, Stephan J.
APPLICANT: Kakavas, Stephan J.
                                                                                                                                                                                                             Query Match
Best Local S
Matches 9
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GENERAL INFORMATION:
                                                                                                        Sequence 32, Patent No. 6
                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, VE
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/718,661
FILING DATE:
                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 530 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ruan, Xiaoan
APPLICANT: Pereda-Lopez, Ana
APPLICANT: Kakavas, Stephan J.
TITLE OF INVENTION: NOVEL POLYKETIDE DERIVATIVES
TITLE OF INVENTION: AND RECOMBINANT METHODS FOR METHODS FOR METHODS FOR METHODS FOR MET
                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 667 amino acids
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                     543 LPVPQPLPQPQMQPQ 557
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US-08-980-459A-2
US-09-583-459A-2
US-09-583-210-2
PCT-US92-00730-2
PCT-US92-00730-2
PCT-US92-00730-2
PCT-US92-00730-2
US-08-804-227C-14
US-08-804-227C-14
US-08-804-108-2
US-08-638-911A-37
US-08-983-359-24
US-08-9893-359-24
US-08-9893-359-24
US-08-9893-358A-2
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CORRESPONDENCE ADDRESS:

ADDRESSEE:

3: Abbott Laboratories 100 Abbott Park Rd.

STREET: 100 Abbott CITY: Abbott Park STATE: Illinois

Illinois

60064-3500

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US-09-078-166-32
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APPLICANT: Katz, I
APPLICANT: Stassi
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                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 49
TELECOMMUNICATION INFORMATION:
TELEPHONE: (847) 938-3137
TELEPHAX: (847)-938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Ruan, Xiaoan
APPLICANT: Pereda-Lopez, Ana
APPLICANT: Kakavas, Stephan J.
TITLE OF INVENTION: NOVEL POLYKETIDE DERIVATIVES
TITLE OF INVENTION: AND RECOMBINANT METHODS FOR PROPERTY OF THE PROPERTY 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
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FILING DATE: 16-MAY-1979
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
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TYPE: amino acid
STRANDEDNESS: sinc
TOPOLOGY: linear
                                                                            OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                        STREET: 100 ADDUCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        42 QLPRPLKDVLFAPEG 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity 46.7%; es 7; Conservation
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REGISTRATION NUMBER: P-40,943
                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
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ARE: FastSEQ Version 2.0 APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2, Application US/09078166
6063561
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                                                                                                                                                                                                                                                                                                                                                                                100 Abbott Park Rd
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Stassi, Diane L.
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                                                                                                        IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                        Abbott Laboratories
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US-08-997-467-32
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                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/8:
FILING DATE: 16-MAX-11997
ATTORNEY/AGENT INFORMATION:
NAME: Dianne Casuto
TELECOMMUNICATION INFORMATION:
TELEPHONE: (847)-938-3137
TELEFAX: (847)-938-2623
                                                                                                                                                                                                                      SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Ruan, Xiaoan
APPLICANT: Pereda-Lopez, Ana
APPLICANT: Kakavas, Stephan J.
TITLE OF INVENTION: NOVEL POLYKETIDE DERIVATIVES
TITLE OF INVENTION: AND RECOMBINANT METHODS FOR MAKING SAME
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REFERENCE/DOCKET NUMBER: 49:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (847)-938-3137
                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
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LENGTH: 343 amino acids
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PRIOR APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                              CITY: Abbott Park STATE: Illinois
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STRANDEDNESS: sin
TOPOLOGY: linear
                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/0 FILING DATE: 16-MAY-1979
                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                       OPERATING SYSTEM:
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Summers Jr., Richard G.
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                                                                                                                                                                                                                                                                       IBM Compatible
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46.78;
                                                                                                                                        08/858,003
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BER: 4952.US.P2
                                                   P-40,943
ER: 4952.US.P2
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; TYPE: PRT
; ORGANISM: Glycine max
US-09-347-833-6
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SEQ ID NO 6
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EARLIER APPLICATION NUMBER: 60/092,415
EARLIER FILING DATE: July 10, 1998
NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/347,833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Famodu, Layo O. APPLICANT: Odell, Joan T.
                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Factors Involved in Gene Expression FILE REFERENCE: BB-1172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 343 amino acids
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                                                                                                                       ADDRESSEE: WOLF, GRE
                                                                                                                                                                          APPLICANT: HELDIN, CARL-HENRIK
TITLE OF INVENTION: PRIMARY STRUCTURE AND FUNCTIONAL
TITLE OF INVENTION: EXPRESSION OF NUCLEOTIDE SEQUENCES FOR NOVEL PROTEIN
TITLE OF INVENTION: TYROSINE PHOSPHATASES
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                         273 QQPMPMMQQQMLPRG 287
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                                       COUNTRY: UZIP: 02210
                                                                                        CITY: BOSTON
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                                                                      MASSACHUSETTS
                                                                                                                                                                                                                                                                                                                                    Application US/08596291
                                                                                                   E: WOLF, GREENFIELD & SACKS, P.C. 600 ATLANTIC AVENUE
                                                                                                                                                                                                                                             CLAESSON-WELSH, LENA
                                                                                                                                                                                                                                                                 GONEZ, LEONEL JORGE SARAS, JAN
                                                      USA
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46.78;
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46.78;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 41; DB
Pred. No. 52;
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RESULT 7
US-09-100-804-4
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Patent No.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 617/720-2441
TELEX: 92-1742 EZEKIEL
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08,
FILING DATE: 01-SEP-1993
ATTORNEY/AGENT INFORMATION:
NAME: GATES, EDWARD R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 09-AUG-19
                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 09-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 617/720-3500
                          APPLICATION NUMBER: US 0: FILING DATE: 01-SEP-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: HELDIN, CARL-HENRIK
TITLE OF INVENTION: EXPRESSION OF NUCLEOTIDE SEQUENCES FOR NOVEL PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                            STREET: BOUL ...
                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            268 KTPDSAQPLLQLMLDVEED 286
                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1
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                                                                                                                                    CLASSIFICATION:
                                                                                                                                                  APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
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REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
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                 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4, Application US/09100804
5. 6066472
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                                                                                                                                                                                                                                                                                                                       MASSACHUSETTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             369 amino acids
                                                                                                                                                                                                                                                                                                                                                        600 ATLANTIC AVENUE
                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GONEZ, LEONEL JORGE SARAS, JAN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PatentIn Release'#1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLAESSON-WELSH, LENA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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01-SEP-1994
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                                                                  US 08/115,573
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                 PCT/US94/09943
                                                                                                US 08/596,291
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Pred. No. 5
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PCT-US94-09943-4
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Best Local :
                                                                     TELEFAX: 617/720-2441
TELEX: 92-1742 EZEKIEL
INFORMATION FOR SEQ ID NO: 4:
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SEQUENCE CHARACTERISTICS:
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TOPOLOGY: line
WOLECUTY
                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                        REFERENCE/DOCKET NUMBER: LO TELECOMMUNICATION INFORMATION: 617/720-3500
                                                                                                                                                                                 APPLICATION NUMBER: US 08
FILING DATE: 01-SEP-1993
ATTORNEY/AGENT INFORMATION:
NAME: TWOMEY, MICHAEL J.
                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Rel
                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
STREET: 600 ATLANTIC AVENUE
                                                                                                                                                                                                                                                  CLASSIFICATION: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE: 01-SE
                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                   TOPOLOGY:
                                           LENGTH:
                                                                                                                                                                       REGISTRATION NUMBER:
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                             : 369 amino acids
amino acid
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TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                USA
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                linear
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47.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYROSINE PHOSPHATASES
                                                                                                                                                                                                                                   US 08/115,573
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Pred. No. 5
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US-08-933-803A-16
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Best Local Similarity
Thes 8; Conserv
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US-09-121-321-16
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Sequence 16, Application US/08933803A Patent No. 6218522 GENERAL INFORMATION:
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Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 212-596-9090 INFORMATION FOR SEQ ID NO: 16:
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 08/
FILING DATE: 19-SEP-1997
ATTORNEY/AGENT INFORMATION:
NAME: Haley, James F
REGISTRATION NUMBER: 27,79
REFERENCE/DOCKET NUMBER: 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Orita, Satoshi
APPLICANT: Igarashi, Hisanaga
APPLICANT: Okumura, Kouichi
APPLICANT: Sakaguchi, Gaku
TITLE OF INVENTION: DNA MOLECULE RELATING TO SUPPRESSION OF
TITLE OF INVENTION: GENE EXPRESSION AND NOVEL PROTEIN
NUMBER OF SEQUENCES: 20
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APPLICANT:
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PRIOR APPLICATION NUMBER: US 08/933,803
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ADDRESSEE: FISH & NEAVE
                                                                                                                415 QPPQPQLQSQPQPQ 428
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/121,321
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                                                                                                                                               2 QTPQPLLQVMMEPQ 15
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57.1%;
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47.4%; Pred. No. 5
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Pred. No. 1
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APPLICANT:
APPLICANT:

Saiga, Akihiko Orita, Satoshi

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Query Match
Best Local Similarity
Thes 8; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 456, Application US/08858207A
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: PATENT: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/933,803A
FILING DATE: 19-SEP-1997
CLASSIFICATION: 536
CCASSIFICATION: 536
                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
ADDRESSEE: SmithKline Beecham Corporation STREET: 709 Swedeland Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Haley, James F.
REGISTRATION UMBER: 27,794
REFERENCE/DOCKET NUMBER: SH
TELECOMMUNICATION INFORMATION:
                                                                                  ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                              APPLICANT: Stodola, Robert TITLE OF INVENTION: No. 6348328el Compounds NUMBER OF SEQUENCES: 552
              COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSQ for Windows Version
CURRENT APPLICATION DATA:
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                                                                                                                                                                         CITY: King of Prussia
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STATE: New York
 APPLICATION NUMBER:
                                                                                                                                         COUNTRY:
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                                                                                                                                         USA
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Knowles, David
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VENTION: DNA MOLECULE RELATING TO SUPPRESSION OF
VENTION: GENE EXPRESSION AND NOVEL PROTEIN
EQUENCES: 20
                                                                                                                                                                                                                                                                                                  Nicholas, Richard
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US/08/858,207A
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Pred. No.
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                                                                         US-09-225-244-2
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                                     Sequence 2, Application US/09225244 Patent No. 6018038
                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 13
LENGTH: 1276
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Best Local
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APPLICANT: PAN, Jae Gu
APPLICANT: PAN, Jae Gu
TITLE OF INVENTION: Membrane-Bound Gluconate Dehydrogenase, Gene Sequence
TITLE OF INVENTION: Encoding the Same and Production of 2-Keto-D-Gluconate
TITLE OF INVENTION: Using Transformed Recombinant E. Coli
FILE REFERENCE: P66159US0
CURRENT APPLICATION NUMBER: US/09/297,937C
CURRENT FILING DATE: 1999-05-11
PRIOR APPLICATION NUMBER: P77/KR98/00296
PRIOR APPLICATION NUMBER: KR 97-48802
PRIOR FILING DATE: 1998-09-25
PRIOR FILING DATE: 1997-09-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13, Application US/09297937C
Patent No. 6337199
APPLICANT: Boung-Jun OH
                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 13
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LENGTH: 179 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 610-270-4478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 60/017670 FILING DATE: 14-MAY-1996 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                              826 LKNPGPLVQAMMK 838
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TOPOLOGY: li
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REGISTRATION NUMBER: 38,891
REFERENCE/DOCKET NUMBER: P50475
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                  INFORMATION:
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Pred. No. 5.
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; TYPE: PRT

; ORGANISM: Capsicum annuum

US-09-225-244-2
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; ORGANISM: Capsicum annuum
US-09-417-242-2
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LENGTH: 328
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Boung-Jun OH
APPLICANT: Moon Kyung KO
APPLICANT: Igor KOSTENYUK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 2
SOFTWARE: Wordperfect 6.1 Windows
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SOFTWARE: Wordperfect 6.1 Windows
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CURRENT FILING DATE: 1999-01-04
EARLIER APPLICATION NUMBER: No. 6018038 Yet Assigned EARLIER FILING DATE: Concurrent Herewith
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APPLICANT: IGOT KOSTENYUK
TITLE OF INVENTION: INCOME
FILE REFERENCE: 1942/35
                 NUMBER OF SEQUENCES: 3

CORRESPONDENCE ADDRESS:

ADDRESSEE: Pretty, Schroeder, Brueggemann & Clark

STREET: 444 South Flower Street, Suite 2000
                                                                                       APPLICANT: MONTMINY, MARC R.
TITLE OF INVENTION: ASSAYS FOR THE IDENTIFICATION OF
TITLE OF INVENTION: COMPOUNDS WHICH INHIBIT ACTIVATION OF CAMP AND MITOGEN
TITLE OF INVENTION: RESPONSIVE GENES
             STREET:
                                                                                                                                                                                                                                                                                                            27 VRKPEPLTQANSDPNG 42
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Los Angeles
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Pred. No. 1
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1.7e+02;
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Search completed: July 31, 2002, 15:23:44 Job time: 147 sec
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Best Local Similarity
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/194,468
FILING DATE: 10-FEB-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                             TELEFAX: (619)-546-9392 INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: P4
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619)-546-4737
                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Reiter, Stephen E.
REGISTRATION NUMBER: 31,192
                                                                                    1902 QTPQPPAQPQPSPVNMSPAG 1921
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LENGTH: 2441 amino aci
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10-FEB-1994
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Pred. No. 1.8e+03;
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seq length: 2000000000
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2: pir2:*
3: pir3:*
4: pir4:*
     July 31, 2002, 15:21:52; Search time 15.13 Seconds (without alignments)
107.966 Million cell updates/sec
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A;Title: The content genome sequence of the lactic acid bacterium Lactococcus lactis A;Reference number: A86625; MUID:21235186; PMID:11337471
A;Accession: D86907
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           elongation factor G [imported] - Lactococcus lactis subsp. lactis (strain IL1403) C;Species: Lactococcus lactis subsp. lactis C;Date: 23-Mar-2001 #sequence_revision 23-Mar-2001 #text_change 03-Aug-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: GB:AE005176; PID:g12725331; A;Experimental source: strain:IL1403 C;Genetics:
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D86907
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C;Superfamily: translation elongation factor G;
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A; Residues: 1-709 <STO>
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translation elongation factor Tu

homo

PIDN:AAK06358.1;

GSPDB:GN00146

translation elonga	EFTWG	_	691	44.9	40	5
esterase, probable	H87344	N	380	44.9	40	14
hypothetical prote	T38405	N	357	44.9	40	ω
protein F1N21.10 [E96696	N	357	44.9	40	42
hypothetical prote	T21149	N	300	44.9	40	Ë
lactoylglutathione	T47277	N	284	44.9	40	ö
hypothetical prote	T14440	N	282	44.9	40	9
hypothetical prote	T14439	N	259	44.9	40	8
hypothetical prote	G72676	N	237	44.9	40	37
protein-tyrosine-p	JC4263	<u>.</u>	656	45.5	40.5	36
protein-tyrosine-p	A55574	_	656	45.5	40.5	5
protein-tyrosine-p	S53427	N	549	45.5	40.5	34
hypothetical prote	F85896	N	1653	46.1	41	ü
hypothetical prote	G65028	N	1653	46.1	41	2
hypothetical prote	в91052	N	1653	46.1	41	Ξ
UL52 protein - hum	WMBE52	Н	1058	46.1	41	30

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A;Gene: lin2571	C;Genetics:	A; Experimental source: strain Clip11262	A;Cross-references: GB:AL592022; PIDN:CAC97798.1; PID:g16415093; GSPDB:GN00178	A; Residues: 1-192 <gla></gla>	A; Molecule type: DNA	A; Status: preliminary	A;Accession: AF1753	A; Reference number: AB1077; MUID:21537279; PMID:11679669	A; Title: Comparative genomics of Listeria species.	ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; Voss, H.; Wehlal	A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; Maitournam, A.,	Science 294, 849-852, 2001	D.; Jones, L.M.; Karst, U.	.; Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussurget, O.; Entian, K.D.; Fsihi,	R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero, F.; Berche, P.; Bloec	C; Accession: AF1753	C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #text_change 27-Nov-2001	C; Species: Listeria innocua	Orf50 [bacteriophage bIL285] homolog lin2571 [imported] - Listeria innocua (strain Cl	AF1753	RESULT 2			15		Qy 1 LQTPOPLLQVMMEPQ 15	Matches 6; Conservative 8; Mismatches 1; Indels 0; Gaps 0;	, 2	

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translation elongation ractor by the state of the state o
                                                                                                                                             A; Gene: SP0273
C; Superfamily:
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C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change
                                                                                                                                                                                                                                  A; Experimental source: strain
                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-693 < KUR>
                                                                                                                                                                                                                                                                                                                                                                               A; Reference number: A95000; A; Accession: B95032
                                                                                                                                                                                                                                                                                                                                                                                                          A; Authors: Loftus, B.J.; Yang, F.; Smith, H.O.; Venter, J.C.; Dougherty, A; Title: Complete Genome Sequence of a virulent isolate of Streptococcus A; Reference number: A95000; MUID:21357209; PMID:11463916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       F;10-17/Region: nucleotide-binding motif A (P-loop)
F;128-131/Region: GTP-binding NKXD motif
F;220-222/Region: GTP-binding SAK/L motif
F;16,17,55,128,129,131,220/Binding site: Mg-GTP (Lys,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R;Sanchez-Pescador, R.; Brown, J.T.; Roberts, M.; Urdea, M.S. Nucleic Acids Res. 16, 1216-1217, 1988
A;Title: The nucleotide sequence of the tetracycline resistance A;Reference number: S03268; MUID:88144009
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                                                                                                                                                                                                                                                             A; Cross-references: GB: AE005672; PIDN: AAK74451.1;
                                                                                                                                                                                                                                                                                                                                                 A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
A; Residues: 1-639 <SAN>
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   h 51.7%;
Similarity 40.0%;
6; Conservative
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Score 46; DB Pred. No. 12; 7; Mismatches
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Pred. No. 1.5;
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                                                                                                                                                elongation
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Radune, D.; Holtzapple,
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A; Gene: fus
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phosphoneuroprotein 14 - bovine
N;Alternate names: brain-specific protein,
C;Species: Bos prinigenius taurus (cattle)
C;Date: 19-May-1994 #sequence_revision 05-M
C;Accession: S39046; JU0238
R;Nakajo, S.; Tsukada, K.; Omata, K.; Nakan
Eur. J. Biochem. 217, 1057-1063, 1993
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Best Local Similarity
"-+-hes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Authors: Yang, Y.; Young-Bellido, M.; Zhao, G.; Zook, C.; Baltz, F. A;Title: Genome of the Bacterium Streptococcus pneumoniae Strain R6. A;Reference number: A97872; MUID:21429245; PMID:11544234 A;Accession: B97903
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C;Accession: B9/700
P. Goskins, J.A.; Alborn Jr.,
                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Title: Identification of two distinct synucleins A;Reference number: $44430; MUID:94252398 A;Accession: $44430 A;Status: preliminary
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R; Jakes, R.; Spillar
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A; Residues: 1-134 < JAK>
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C;Date: 07-Oct-1994 #sequence_revision 10-Nov-1995
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A; Residues: 1-693 <KI
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;Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 02-Nov-2001
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A;Authors: Hunter, J.L.; Jenkins, J.; Johnson-Hopson, C.; Khan, S.; Khan, C.A.; Li, J.H.; Li, Y.; Lin, X.; Liu, S.X.; Liu, Z.A.; Luros, J.S.; Mail Rizzo, M.; Rooney, T.; Rowley, D.; Sakano, H.

A;Authors: Salzberg, S.L.; Schwartz, J.R.; Shinn, P.; Southwick, A.M.; & Ker, M.; Wu, D.; Yu, G.; Fraser, C.M.; Venter, J.C.; Davis, R.W.

A;Title: Sequence and analysis of Chromosome 1 of the plant Arabidopsis A;Reference number: A86141; MUID:21016719
                                                                                                                                                                                                                                           C;Date: 02-Mar-2001 #sequence_revision C;Accession: C96534
R;Theologis, A.; Ecker, J.R.; Palm, C.J. Chin, C.W.; Chung, M.K.; Conn, L.; Con ansen, N.F.; Hughes, B.; Hulzar, L. Nature 408, 816-820, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     R;Tobe, T.; Nakajo, S.; raman, ...
J. Neurochem. 59, 1624-1629, 1992
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C;Species: Rattus sp. (rat)
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 05-Nov-1999
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-671 <STO>
A;Cross-references: GB:AE005173; NID:g10120431; PIDN:AAG13056.1; GSPDB:GN00141
                                                                                       A; Reference number: A86141; A; Accession: C96534
                                                                                                                                                                                                                                                                                                                                         probable Poly-A Binding Protein [imported] - Arabidopsis thaliana C;Species: Arabidopsis thaliana (mouse-ear cress) C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 23-Mar-2001
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C; Keywords: brain
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A; Accession: JU0238
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Residues: 1-134 <NA2>
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                                                                                                                                                                                                                                                                                   C.J.; Federspiel, N.A.; Kaul, S.; W Conway, A.B.; Conway, A.R.; Creasy,
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Maiti, R.; M
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A;Title: Nucleotide sequence of the tet(M)
A;Reference number: S13142; MUID:91045089
A;Accession: S13142
C;Superfamily: translation elongation factor G; translation elongation factor C;Eeywords: antibiotic resistance; GTP binding; nucleotide binding; P-loop F;4-131/Domain: translation elongation factor Tu homology <ETU>F;10-17/Region: nucleotide-binding motif A (P-loop)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nature 413, 848-852, 2001
A;Authors: Parry, C.; Quail, M.; Rutherford, K.; Simmonds, M.;
A;Title: Complete genome sequence of a multiple drug resistant
                                                                                                                                                                                                                                                                                                                                                             C; Species: Enterococ
C; Date: 13-Jan-1995
C; Accession: S13142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    conserved hypothetical protein STY2446 [imported] - Salmonel C; Species: Salmonella enterica subsp. enterica serovar Typhi A; Note: this species has also been called Salmonella typhi
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AB0784
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C;Superfamily: polyadenylate-binding protein;
                                                                     A; Gene:
                                                                                                         A; Contents:
A; Note: pro
                                                                                                                        A; Reference number: A23749; MUID:91131580 A; Contents: annotation; function; amino end
                                                                                                                                                        J. Biol. Chem. 266, 2872-2877, 1991
A; Title: Purification and characterization of Tet(M), a
                                                                                                                                                                                                                                  A; Molecule type: DNA
A; Residues: 1-639 <BUR>
                                                                                                                                                                                                                                                                                                                                                                                               tetracycline resistance protein tetM -
C;Species: Enterococcus faecalis
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A; Residues: 1-134 < PAR>
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                                                                                        C; Genetics
                                                                                                                                                                                                R;Burdett,
                                                                                                                                                                                                                 A;Cross-references: EMBL:X56353; NID:g47061; PIDN:CAA39796.1; PID:g47062
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8; Conser
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A;Residues: 1-639 <SU1>
A;Cross-references: GB:M85225; NID:g148321; PIDN:AAA24784.1; PID:g148322
A;Cross-references: GB:M85225; NID:g148321; PIDN:AAA24784.1; PID:g148322
A;Note: sequence extracted from NCBI backbone (NCBIN:111076)
C;Superfamily: translation elongation factor G; translation elongation factor C;Keywords: antibiotic resistance; GTP binding; nucleotide binding; P-loop F;4-131/Domain: translation elongation factor Tu homology <ETU>
F;10-17/Region: nucleotide-binding motif A (P-loop)
F;128-131/Region: GTP-binding NKXD motif F;220-222/Region: GTP-binding SAK/L motif
F;20-222/Region: GTP-binding SAK/L motif
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A;Recession: A60633
A;Molecule type: DNA
A;Residues: 1-639 <NES>
A;Residues: 1-639 <NES>
A;Cross-references: GB:M21136; NID:9153114; PIDN:AAA26678.1; PID:9153115
C;Superfamily: translation elongation factor G; translation elongation fact
C;Keywords: antibiotic resistance; GTP binding; nucleotide binding; P-loop
F;4-131/Domain: translation elongation factor Tu homology <ETU>
F;10-17/Region: nucleotide-binding motif A (P-loop)
F;10-17/Region: GTP-binding SAK/L motif
F;220-222/Region: GTP-binding SAK/L motif
F;128-131/Tegion: GTP-binding SAK/L motif
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                                                                                                                                                                                                                                                                                                                                                                                                                                    cetracycline resistance protein TetM - Enterococcus faecalis tra
C;Species: Enterococcus faecalis
C;Date: 08-Sep-1995 #sequence_revision 08-Sep-1995 #text_change
C;Accession: A56779
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C; Accession:
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F;220-222/Region: GTP-binding SAK/L motif
F;16,17,55,128,129,131,220/Binding site: Mg-GTP
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    Similarity
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    Score
Pred.
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D.; Jones, L.M.; Karst, U. Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-EA;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Accession: AE1406
                                                                                                                                                                                                                                                                             D.; Jones, L.M.; Karst, U. Science 294, 849-852, 2001
A;Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Maduenc ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, A;Title: Comparative genomics of Listeria species.
A;Reference number: AB1077; MUID:21537279; PMID:11679669
                                                                                                                                                                                                                                                                                                                                                                                   C;Accession: AE1782
R;Glaser, P.; Frangeul, L.; Buchrieser, C.; A.
.; Dominguez-Bernal, G.; Duchaud, E.; Durand,
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C;Accession: AE1406
R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.; Baquero,
                                                                                                                       A; Gene: fus
C; Superfamily:
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C; Date: 27-Nov-2001
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A; Residues: 1-695 <G
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A;Experimental source: strain Clip11262
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Pfam; PF00679; EFG_C; 1.

Pfam; PF00009; GTP_EFTU; 1.

Pfam; PF03144; GTP_EFTU_D2; 1.

PRINTS; PR00315; ELONGATNFCT.

PROSITE; PS00301; EFACTOR_GTD; 1.

Elongation factor; Protein biosynthesis; GTP-binding;
                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation—the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                      EMBL; AE006493; AAK33347.1; -.
InterPro; IPR000640; EFG_C.
InterPro; IPR000795; GTP_EFTU.
InterPro; IPR004161; GTP_EFTU_D2.
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Result No.

3328876544 3328876544 3328876544 3328876544

Database :

ALIGNMENTS

EFG_STRPY STANDARD; PRT; 691 AA.
16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
(Rel. 40,
FUS OR SPY0273.
Streptococcus pyogenes.
Bacteria; Firmicutes; Bacillus/Clostridium group; Streptococcaceae;
Streptococcus.
SEQUENCE FROM N.A.
MEDLINE=21192684; PubMed=11296296;
Ferretti J.J., McShan W.M., Ajdic D.J., Savic D.J., Savic G., Lyon K.,
Qian Y., Jia H.G., Najar F.Z., Ren Q., Zhu H., Song L., White J.,
Yuan X., Clifton S.W., Roe B.A., McLaughlin R.;
Proc. Nath Acad. Sci. II.S.A. 98:4658-4663(2001).
PARTIAL SEQUENCE, AND MASS SPECTROMETRY.
STRAIN-JRS4 / Serotype M6;
Hogan D.A., Du P., Stevenson T.I., Whitton M., Kilby G.W., Rogers J.,
"Two-dimensional gel electrophoresis map of Streptococcus pyogenes
proteins.";
Submitted (MAY-2000) to the SWISS-PROT data bank!- FUNCTION: THIS PROTEIN PROMOTES THE GTP-DEPENDENT TRANSLOCATION
OF THE NASCENT PROTEIN CHAIN FROM THE A-SITE TO THE P-SITE OF THE
RIBOSOME (BY SIMILARITY)I- SUBCELLULAR LOCATION: Cytoplasmic.
-!- SIMILARITY: BELONGS TO THE GTP-BINDING ELONGATION FACTOR FAMILY.EF-G/EF-2 SUBFAMILY.
This culture books is converted. It is produced through a collaboration

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PIR; S03268; S03268.
HSSP; P13551; 1FNM.
InterPro; IPR000640; EFG_C.
InterPro; IPR000795; GTP_EFTU_InterPro; IPR0004161; GTP_EFTU_D2.
Pfam; PF00679; EFG_C; IPFfam; PF00009; GTP_EFTU; 1.
Pfam; PF00009; GTP_EFTU, D2; 1.
PF1MT; PF00144; GTP_EFTU_D2; 1.
                                                                                                                                                  NP_BIND
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INIT_MET
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=88144009; PubMed=3344217;
Sanchez-Pescador R., Brown J.T., Roberts M., Urdea M.S.;
Sanchez-Pescador R., Brown J.T., Roberts M., Urdea M.S.;
"The nucleotide sequence of the tetracycline resistance determinant tetm from Ureaplasma urealyticum.";
Nucleic Acids Res. 16:1216-1217(1988).
                                                                                                                                                                    Protein biosynthesis; Antibiotic NP_BIND 10 17 GTP NP_BIND 74 78 GTP NP_BIND 128 131 GTP
                                                                                                                                                                                                                                                         PRINTS; PR00315; ELONGATNFCT. PROSITE; PS00301; EFACTOR_GTP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
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01-MAR-1989 (Rel.
16-OCT-2001 (Rel.
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Bacteria; Firmicutes; Bacillus/Clostridium group; Mollicutes;
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Pred. No. 3.8;
4; Mismatches
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GTP (BY SIMILARITY).

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A1497055BB182B3A CR
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D59B857A2CDB40CD CR
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                                                   Query Match
Best Local
                                         Matches
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P33567;
01-FEB-1994
                                                                                                                                                                                                  Pfam; PF01387; Synuclein; 1
PRINTS; PR01211; SYNUCLEIN.
ProDom; PD010631; Synucleir
                                                                                                                                                                                                                                                                                                          J. Neurochem. 55:2031-2038(1990).
-i- FUNCTION: MAY BE INVOLVED IN NEURONAL
-i- SUBCELLULAR LOCATION: CYTOPLASMIC.
-i- TISSUE SPECIFICITY: SPECIFICALLY PRESE
NEURONS BUT NOT IN GLIAL CELLS.
-i- PTM: PHOSPHORYLATION B
                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=94039126; pubMed=8223629;
Makajo S., Tsukada K., Omata K., Nakamura
"A new brain-specific 14-kDa protein is a
complete amino acid sequence and evidence
Eur. J. Blochem. 217:1057-1063(1993).
                                                                                                                REPEAT
MOD_RES
                                                                                                                                     REPEAT
                                                                                                                                                                                                                                            PIR; JU0238; JU0238.
PIR; S39046; S39046.
                                                                                                                                                                                                                                                                 -!- SIMILARITY: BELONGS
PIR; A60218; A60218.
                                                                                            SEQUENCE
                                                                                                                                                                                          Phosphorylation;
                                                                                                                                                                                                                                                                                                                                                                         protein."
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Nakai Y., Nakaya K., Nakamura Y.;
                                                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Brain;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pec
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specific protei
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01-FEB-1994 (Rel.
16-OCT-2001 (Rel.
                                                                                                                                                  REPEAT
                                                                                                                                                           REPEAT
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                                                                                                                                                                                                                                    InterPro; IPR001058;
                                                                                                                                                                                                                                                                                                                                                                                   "Purification and characterization
                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=91038084; PubMed=2230807;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bovidae; Bovinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bos taurus
105
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EPLIEPLMEPEGE
                  OPLLOVMMEPOGD
                                        6; Conserv
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118
                                         Conservative
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el. 28, Last sequence update)
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(Phosphoneuroprotein 14) (PNP
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67
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                                                                                                                                                                                                                                    Synuclein
                                                    49
                                                   . 28;
                                                                                                                                                                                                                                                                             TO THE SYNUCLEIN FAMILY
                                                                                             MW.
                                         6
                                        Score 44; DB Pred. No. 1.5; 6; Mismatches
                                                                                                       (BY
                                                                                                                PHOSPHORYLATION (BY CK2,
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[EQ]-[GQ]-V-X(4).
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phosphoprotein. Its
for phosphorylation.";
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                                                             Length 134;
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Pecora; Bovoidea
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RESULT

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SYUB_HUMAN

SYUB_HIMAN

OXUB_H

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16-OCT

DE Beta-s

GNCB.
SNCB.
OS HOMO S

OC EUKARY

OC MAMMADIA

NOSH_T

RN [1]

RP SEQUEN

RA JAKES

RT TAKES

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01-NOV-1997
01-NOV-1997
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-I- FUNCTION: MAY BE INVOLVED IN NEURONAL PLASTICITY.
-I- SUBCELLULAR LOCATION: Cytoplasmic.
-I- TISSUE SPECIFICITY: EXPRESSED PREDOMINANTLY IN BRAIN; CONCENTRATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; C
Mammalia; Eutheria; P
       REPEAT
REPEAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pronin A.N., Morris A.J., Surguchov A., Benovic J.L.;
"Synucleins are a novel class of substrates for G protein-coupled receptor kinases.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Jakes R., Spillantini M.G., Goedert M.; "Identification of two distinct synucleins from human brain."; FEBS Lett. 345:27-32(1994).
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01-NOV-1997 (Rel. 35, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
MEDLINE=99026142; PubMed=9806846;
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                                                                                                                                                                          Phosphorylation;
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ProDom; PD010631;
                                                                                                                                                                                                                                                                          PRINTS;
                                                                                                                                                                                                                                                                                                                                                                                    MIM; 602569;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genomics 54:173-175(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=94252398; PubMed=8194594;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE-Brain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID-9606;
                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR001058; Synuclein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SNCB)."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AND CAM-KINASE II.

DISEASE: BRAIN IRON ACCUMULTION TYPE 1 (NBIA1, ALSO CALLED DISEASE: BRAIN TRON ACCUMULTION TYPE 1 (NBIA1, ALSO CALLED HALLERVORDEN-SPATZ SYNDROME), A RARE NEUROAXONAL DYSTROPHY, IS HISTOLOGICALLY CHARACTERIZED BY AXONAL SPHEROIDS, IRON DEPOSITION, LEWY BODY (LB)-LIKE INTRANEURONAL INCLUSIONS, GLIAL INCLUSIONS AND LEWY BODY (LB)-LIKE INTRANEURONAL INCLUSIONS INCLUSIONS AND LEWY BODY (LB)-LIKE INTRANEURO (LB)-LIKE 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IN PRESYNAPTIC NERVE TERMINALS.
PTM: PHOSPHORYLATED. PHOSPHORYLATION
KINASES (GKR) IS MORE EFFICIENT THAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY: BELONGS TO THE SYNUCLEIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NEUROFIBRILLARY TANGLES. SNCB IS FOUND IN SPHEROIDS BUT NOT IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SWISS-PROT entry is copyright. It is produced through a collaboration een the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INCLUSIONS.
                                                                                                                                                                                                                                                                                                                                                                                                             $69965; AAB30860.1; -. AF053136; AAC80286.1; -. AF053134; AAC80286.1; JC AF053135; AAC80286.1; JC
                                                                                                                                                                                                                                                                      s; PR01211; SYNUCLEIN: 1
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                                                                                                                                                                          Repeat.
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                                                                                                                                                                                                                                             Synuclein; 1.
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Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AND
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                                                                                                                                                                                                                                                                                                                                                                                                                 JOINED.
                                                                                                                                      4 x 11 AA TANDEM [EQ]-{GQ}-V-X(4).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRT;
                                   (APPROXIMATE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BY G-PROTEIN COUPLED RE PHOSPHORYLATION BY CK1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      human beta-synuclein gene
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                                                                                                                                                                          REPEATS
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                                                                                                                                                                          [EGS]-K-T:K-
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SYUB_RAT
Q63754;
                                                                                                                                                                                                                                                                                                      Nakajo S., Tsukada K., Omata K., Nakamura Y., Nakaya K.;

"A new brain-specific 14-kDa protein is a phosphoprotein. Its complete
amino acid sequence and evidence for phosphorylation.";

Eur. J. Biochem. 217:1057-1063(1993).

-!- FUNCTION: MAY BE INVOLVED IN NEURONAL PLASTICITY.

-!- TISSUE SPECIFICITY: EXPRESSED SPECIFICALLY IN BRAIN.

-!- PTM: PHOSPHORYLATED. PHOSPHORYLATION BY G-PROTEIN COUPLED RECEPTOR

KINASES (GKR) IS MORE EFFICIENT THAN PHOSPHORYLATION BY CK1, CK2

AND CAM-KINASE II.

-!- SIMILARITY: BELONGS TO THE SYNUCLEIN FAMILY.
                                                             REPEAT
                                                                                                                                                                                                  use by non-profit institutions as long modified and this statement is not removed. entities requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entropean Bioinformatics Institute. There are no resture by non-profit institutions as long as its content
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Cloning and characterization of specific 14-kDa protein."; J. Neurochem. 59:1624-1629(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-OCT-2001 (Rel. 40, Last annotation update)
Beta-synuclein (Phosphoneuroprotein 14) (PNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation updat
 SEQUENCE
                                                                                                                          PRINTS; PR01211; S
ProDom; PD010631;
                                                                                                                                                  InterPro; IPR001058; Synuclein.
Pfam; PF01387; Synuclein; 1.
                                                                                                                                                                              EMBL; D17764; BAA04610.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.,
                                                                           REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=94039126; PubMed=8223629;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tomita M.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=93019240;
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                       MOD_RES
                                                  REPEAT
                                                                                                              Phosphorylation;
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                                                                                                  DOMAIN
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6; Conser
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134 AA;
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 137
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                                                                                                  Repeat.
67
                                                                                                                                      SYNUCLEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PubMed=1402909,
                                                                                                                           Synuclein;
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 14504
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Pred. No.
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; 5BCA9FCA615AC4EF CRC64;
(BY SIMILARITY). 678C6CB84FA01A03 CRC64;
                      PHOSPHORYLATION (BY CK2,
                                                                                     4 X 11 AA TANDEM REPEATS [EQ]-[GQ]-V-X(4).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Craniata; Vertebrata; Euteleostomi;
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                                                 (APPROXIMATE)
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                                                                                                                                                                                                               moved. Usage by and for (See http://www.isb-sib.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   encoding a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Omata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             K., Nakaya K.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    novel
                          GRK2 AND
                                                                                                   OF [EGS]-K-T-K-
                                                                                                                                                                                                                                                      restrictions
                                                                                                                                                                                                                                                                    EMBL
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                                                                                                                                                                                                                                                                    outstation
                          GRK5)
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Best Local
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InterPro; IPR000795; GTP_EFTU.
InterPro; IPR004101; GTP_EFTU_D2.
Pfam; PF00679; EFG_C; 1.
Pfam; PF00009; GTP_EFTU, 1.
Pfam; PF03144; GTP_EFTU_D2; 1.
                                                                                                                                                                                                           Transposable
NP_BIND
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NP_BIND
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SEQUENCE 63
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TET9_ENTFA
P21598;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation—the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; x56353; CAA39796.1;
PIR; S13142; S13142.
HSSP; P13551; 1FNM.
                                                                                                                                                                                                                                                                                                                            PRINTS; PR00315; ELONGATHFCT.
PROSITE; PS00301; EFACTOR_GTP; 1.
Protein biosynthesis; Antibiotic resistance;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAY-1991 (Rel. 18, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Tetracycline resistance protein tetM from transposon Tn916
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Nucleotide sequence of the tet(M) gene of Tn916."; Nucleic Acids Res. 18:6137-6137(1990).
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TETM OR TET(M).
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01-MAY-1991 (Rel.
16-OCT-2001 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Burdett V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=91045089; PubMed=2172929;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=1351;
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Bacteria; Firmicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-MAY-1991
339
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                                                                                                                Local
                                            1 LQTPQPLLQVMMEP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FUNCTION: ABOLISH THE INHIBITORY EFFECT OF TETRACYCLIN ON PROTEIN SYNTHESIS BY A NON-COVALENT MODIFICATION OF THE RIBOSOMES.
SIMILARITY: BELONGS TO THE GTP-BINDING ELONGATION FACTOR FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TETM/TETO SUBFAMILY.
IENPHPLLQTTVEP
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                                                                                          Similarity
7; Conser
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                                                                                                                                                                                                             128
639 /
                                                                                             Conservative
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Bacillus/Clostridium group;
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Pred. No. 1.5;
6; Mismatches
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                                                                                                                                                                                                        GTP (BY SIMILARITY).
GTP (BY SIMILARITY).
GTP (BY SIMILARITY).
; B2315A37B53B18FB CRC64;
                                                                                                              Score 43; DB
Pred. No. 12;
                                                                                             Mismatches
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                                                                                                                                    Length 639;
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Best Local :
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yfam; PF0314; GTP_EFTU___,

PRINTS; PR00315; ELONGATNFCT.

PROSITE; PS00301; EFACTOR_GTP; 1.

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Cloning and nucleotide sequence of a chromosomally encoded tetracycline resistance determinant, tetk(M), from a pathogenimethicillin-resistant strain of Staphylococcus aureus.";
Antimicrob. Agents Chemother. 34:2273-2276(1990).
- FUNCTION: ABOLISH THE INIBITORY EFFECT OF TETRACYCLIN ON SYNTHESIS BY A NON-COVALENT MODIFICATION OF THE RIBOSOMES.
-I- SUMILARITY: BELONGS TO THE GTP-BINDING ELONGATION FACTOR F
                                Bacteria; Proteobacteria; Rhizobiaceae; Rhizobium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; M21136; AAA26678.1; -. HSSP; P13551; 1FNM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
MEDLINE-91158314; PubMed-2073121;
Nesin M., Svec P., Lupski J.R., G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tetracycline resistance TETM OR TETA(M).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-DEC-1998
15-DEC-1998
15-DEC-1998
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InterPro; IPR000795;
InterPro; IPR004161;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entities requires a license agreement (See http://www.isb-sib.ch/announce/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TETM_STAAU
Q53770;
                                                                                                         Agrobacterium tumefaciens
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PF00009; GTP_EFTU; 1.
PF03144; GTP_EFTU_D2; 1.
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                                                                     group;
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Best Local
                                                                                                                                                                                                                                                                                                                               P52369;
P52369
SEQUENCE FROM N.A.

MEDLIND-95302501; PubMed-7783207;

Telford E.A., Watson M.S., Aird H.C., Perry J., Davison A.J.;

Telford E.A., Watson M.S., Aird H.C., Perry J., Davison A.J.;

The DNA sequence of equine herpesvirus 2.";

J. Mol. Biol. 249:520-528(1995)

-!- FUNCTION: THE CAPSID ASSEMBLY PROTEIN IS COMPONENT OF THE CAPSID CORE INVOLVED IN PROCESSING AND PACKAGING OF PROGENY DNA.

-!- FUNCTION: ASSEMBLIN IS A PROTEASE WHICH CAN PROTEOLYTICALLY CLEAVE ITSELF AND THE CAPSID ASSEMBLY PROTEIN AT THE C-TERMINUS.

-!- CATALYTIC ACTIVITY: PREFERENTIALLY CLEAVES AT ALA-|-SER OR ALA-|-

ALA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NP_BIND
NP_BIND
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SEQUENCE
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PROSITE;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and fo entities requires a license agreement (See http://www.isb-sib.
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MEDLINE=97047977; PubMed=8892818;
Syvanen A., Amiri H., Jamel A., A
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                                                                                                                                                                                                                                                           Gammaherpesvirinae.
NCBI_TaxID=82831;
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                                                                                                                                                                                                                                                                                                           Viruses; dsDNA viruses,
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InterPro;
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rPro; IPR000795; GTP_EFTU_D2.
rPro; IPR004161; GTP_EFTU_D2.
n; PF00679; EFG_C; 1.
n; PF000099; GTP_EFTU_D2; 1.
n; PF003144; GTP_EFTU_D2; 1.
site; PS00301; EFACTOR_GTP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          European Bioinformatics Institute.
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SIMILARITY: BELONGS TO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EF-G/EF-2 SUBFAMILY.
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Best Local
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STRAIN-KIZ / MG1655;

MEDLINE-97426617; pubMed=9278503;

MEDLINE-97426617; pubMed=9278503;

Blattner F.R., Plunkett G. III, Bloch C.A., Perna N.T., Burl Blattner F.R., Plunkett G. III, Bloch C.A., Rode C.K., Mayhew Riley M., Collado-Vides J., Glasner J.D., Rode C.K., Mayhew Riley M., Collado-Vides J., Glasner J.D., Goeden M.A., Rose I Davis N.W., Kirkpatrick H.A., Goeden M.A., Rose I
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                                    Perna N.T., Plunkett G. III, Burland V., Mau B., Glasner J.D., Rose D.J., Mayhew G.F., Evans P.S., Gregor J., Kirkpatrick H.A., Posfai G., Hackett J., Klink S., Boutin A., Shao Y., Miller L., Grotbeck E.J., Davis N.W., Lim A., Dimalanta E.T., Potamousis K., Apodaca J., Anantharaman T.S., Lin J., Yen G., Schwartz D.C., Welch R.A., Blattner F.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      use by non-profit institutions as long modified and this statement is not removed entitles requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                                                                                SEQUENCE FROM N.A. STRAIN-0157:H7 / E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hypothetical protein yhgG. YHGG OR B3410 OR Z4765 OR Escherichia coli, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; U20824; AAC13804.1; -. HSSP; P16753; 1CMV.
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MEDLINE=21074935; PubMed=11206551;
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SIMILARITY: BELONGS TO PEDTIDASE FAMILY S21 (SERINE PROSIMILARITY: TO OTHER HERPESVIRUSES CAPSID PROTEIN VP40.
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                                                                                                                                                                                                                                                                complete genome sequence nce 277:1453-1474(1997).
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SEQUENCE FROM N
STRAIN=0157:H7 ,
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01-FEB-1996 (Rel. 33, Created)
01-OCT-1996 (Rel. 34, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
T-complex protein 1, gamma subunit (TCP-1-gamma) (CCT-gamma).
CCT-GAMMA OR CCT-3 OR CCTG OR CGB977.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=0157:H7 / RIMD 0509952;

MEDLINE=21156231; PubMed=11258796;

Hayashi T., Makino K., Ohnishi M., Kurokawa K., Ishii K., Yokoya Hayashi T., Makino E., Nakayama K., Murata T., Tanaka M., Tobe Tida T., Takami H., Honda T., Sasakawa C., Ogasawara N., Yasunag Kuhara S., Shiba T., Hattori M., Shinagawa H.;

"Complete genome sequence of enterohemorrhagic Escherichia coli 0157:H7 and genomic comparison with a laboratory strain K-12.";

DNA Res. 8:11-22(2001).
  Adams M.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., Amanatides P.G., Scherer S.E., Li P.W., Hoskins R.A., Galle R.F., George R.A., Lewis S.E., Richards S., Ashburner M., Henderson S.N., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X.,
                                                                                                         STRAIN-BERKELEY;
MEDLINE-20196006; PubMed-10731132;
                                                                                                                                                                                                                        "Drosophila melanogaster P1 geno chaperonin-encoding gene Cctg."; Gene 171:221-223(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Musc
Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EcoGene; EG12933; yhgG.
Hypothetical protein; Complete proteome.
SEQUENCE 78 AA; 8660 MW; 88976DE22CA9024B CRC64;
                                                                                                                                                                                                                                                                                                       Walkley N.A., Malik A.N.;
                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                      Submitted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DROME
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                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                 MEDLINE-96257223;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRAIN-CANTON-S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=7227;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26
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D.R., Celniker S.E
ted (JUL-1995) to ti
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46.28;
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., Lewis E.B., Palazzolo M.J.;
he EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                           genomic
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Pred. No. 2.
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RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA Cherry J.M., Cawley S., Dahlke C., Ferraz C., Ferriera S., Fleischmann W.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Fosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Melson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Mount S.M., Welson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Mullians S.M., Welson K.A., Weissenbach J.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,
RA Kinnes S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,
RA Kinnes S.M., Woodage T., Worley K.C., Wu D., Yang S., Zhou Q., Zheng L.,
RA Chilam S.M., Woodage T., Worley K.C., Wang A.H.,
                              Query Match
Best Local
                                                                                                                                                           Chaperone;
CONFLICT
CONFLICT
SEQUENCE
                                                                                                                                                                                                                                                                                 PROSITE; PS00750; TCP1_1; 1.
PROSITE; PS00751; TCP1_2; 1.
PROSITE; PS00995; TCP1_3; 1.
                                                                                                                                                                                                                                                                                                                                                                                      Interpro; IPR002423; TCP1_cpn6
Interpro; IPR002194; TCP_1
Pfam; PF00118; cpn60_TCP1; 1.
PRINTS; PR00304; TCOMPLEXTCP1.
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EMBL; X95602; CAA64860.1; --
EMBL; AE003714; AAF55350.1;
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FlyBase; FBgn0015019; Cct-gamma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBCELLULAR LOCATION: Cytoplasmic.
SIMILARITY: BELONGS TO THE TCP-1 CHAPERONIN FAMILY.
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SUBUNIT: HETERO-CLIGOMERIC COMPLEX OF ABOUT 850 TFORMS TWO STACKED RINGS, 12 TO 16 NM IN DIAMETER.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FUNCTION: MOLECULAR CHAPERONE; ASSIST THE FOLDING ATP HYDROLYSIS. KNOWN TO PLAY A ROLE, IN VITRO, II
                                                                                                                                                12
35
544
                                                                                                                                                           ATP-binding; Multigene family.
12 S -> SD (IN REF. 2).
35 47 MISSING (IN REF. 1).
544 AA; 59394 MW; DD5E635809B322C2 C
                          46.18;
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Score 41; DB
Pred. No. 22;
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01-JUL-1989 (Rel. 11, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              TET5_ENT P11131;
                                          STRPN
                                                                                                                                                                                                                       PRINTS; PR00315; ELONGATNECT.
PROSITE; PS00301; EFACTOR_GTP; 1.
Protein biosynthesis; Antibiotic resistance;
                                                                                                                                                                                                                                                     Pfam; PF00679; EFG_C; 1.
Pfam; PF00009; GTP_EFTU, 1.
Pfam; PF03144; GTP_EFTU_D2; 1.
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                                                                                                                                                                                                                Transposable element
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InterPro; IPR000795; GTP_EFTU.
InterPro; IPR004161; GTP_EFTU_D2.
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128
639
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                                                                                                                       Conservative
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50.0%;
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Bacillus/Clostridium group;
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                                                                                                                       Score 41; DB
Pred. No. 26;
3; Mismatches
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GTP (BY SIMILARITY).
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                                                                                                                                           Length 639;
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Q48791;
15-DEC-1998
15-DEC-1998
15-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                             NP_BIND
NP_BIND
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEMS Microbiol. Lett. 135:231-236(1996).

-I- FUNCTION: ABOLISH THE INHIBITORY EFFECT OF TETRACYCLIN ON PROTEI SYNTHESIS BY A NON-COVALENT MODIFICATION OF THE RIBOSOMES.
-I- SIMILARITY: BELONGS TO THE GTP-BINDING ELONGATION FACTOR FAMILY.

TETM/TETO SUBFAMILY.
                                                              Plasmid pIP811.

Bacteria; Firmicutes; Bacillus/Staphylococcus
                                                                                                                   Tetracycline resistance TETS OR TET(S).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00679; EFG_C; 1.
Pfam; PF00009; GTP_EFTU; 1.
Pfam; PF03144; GTP_EFTU_D2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; X90939; CAA62436.1; -. HSSP; P13551; 1FNM. InterPro; IPR000640; EFG_C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIINE=96162866; PubMed=8595862;
Provvedi R., Manganelli R., Pozzi G.;
"Characterization of conjugative transposon Tn5251 of Streptococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Streptococcus pneumoniae.
Bacteria; Firmicutes; Bacillus/Clostridium group; Streptococcaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-DEC-1998 (Rel. 37, Last annotation update)
Tetracycline resistance protein tetM from transposon
TETM(5251).
              SEQUENCE FROM N.A STRAIN=BM4210;
                                                                                                                                                                                                                                                                                                                                                                                                                   Transposable
NP_BIND
                                                                                                                                                                                                                                                                                                                                                                                                                                            PROSITE; PS00301; EFACTOR_GTP; 1. Protein biosynthesis; Antibiotic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Interpro; IPR000795; GTP_EFTU.
Interpro; IPR004161; GTP_EFTU_D2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pneumoniae.",
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
MEDLINE=93380670; PubMed=8370538;
                                                                                                     Listeria monocytogenes.
                                                                                                                                                                                                            LISMO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=1313;
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639 4
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(Rel. 37, Last sequence update)
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72556 MW;
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50.0%;
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78
                                                                              Bacillus/Clostridium
                                                                                                                              protein tetS (Tet(S))
                                                               group;
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                                                                                                                                                                                                                                                                                                                                                                                           GTP
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                                                                                                                                                                                                PRT;
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TP (BY SIMILARITY).
F18131E08B81F3C0 CRC64;
                                                                 Listeria
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Matches 7
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InterPro; IPRO00795; GTP_EFTU.
InterPro; IPRO04161; GTP_EFTU_D2.
Pfam; PF00679; EFG_C; 1.
Pfam; PF00009; GTP_EFTU_D2; 1.
Pfam; PF031444; GTP_EFTU_D2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TETS_LACLA STAN
Q48712;
15-DEC-1998 (Rel. 3
15-DEC-1998 (Rel. 3
16-OCT-2001 (Rel. 4
                                                                                                                                                                           STRAIN-K214;
Strain-K214;
Perreten V., Schwarz F., Cresta L., Boeglin M., Dasen G., Teuber M.;
"Antiblotic resistance spread in food.";
"Antiblotic resistance spread in food.";
Submitted (OCT-1997) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: ABOLISH THE INHIBITORY EFFECT OF TETRACYCLIN ON PROTEIN SYNTHESIS BY A NON-COVALENT MODIFICATION OF THE RIBOSOMES.
-I- SIMILARITY: BELLONGS TO THE GTP-BINDING ELONGATION FACTOR FAMILY.
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                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TETS OR TET(S)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Plasmid pK214.
Bacteria; Firmicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRINTS; PR00315; ELONGATNFCT.
PROSITE; PS00301; EFACTOR_GTP; 1.
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"Characterization of a new class of tetracycline-resistance tet(s) in Listeria monocytogenes BM4210.";
Gene 131:27-34(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID-1360;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lactococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tetracycline resistance
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                                                                                                                                                                  TETM/TETO SUBFAMILY.
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SIMILARITY: BELONGS TO THE GTP-BINDING ELONGATION FACTOR FAMILY.
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74 78 GTP
128 131 GTP
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131 6
; 73014 MW;
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ce protein tetS (Tet(S
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Pred. No. 26;
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GTP (BY SIMILARITY).
GTP (BY SIMILARITY).
GTP (BY SIMILARITY).
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(Tet(S)).
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Matches 7
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NP_BIND
SEQUENCE
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Pfam; PF00009; GTP_EFTU; 1.
Pfam; PF03144; GTP_EFTU_D2; 1.
                                                                                                                                                               PRINTS; PR00315; ELONGATNFCT.
PROSITE; PS00301; EFACTOR_GTP; 1.
                                                                                                                                                                                                                      InterPro; IPR000640; EFG_C.
InterPro; IPR000795; GTP_EFTU.
InterPro; IPR004161; GTP_EFTU_D2.
                                                                                                                                                                                                                                                         EMBL; X92946; CAA63528.1; -. HSSP; P13551; 1FNM.
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                                                                                                                                        NP_BIND
                                                                                                                                                     Protein
344 LENPLPMLQTTIEP
                       Н
            LQTPQPLLQVMMEP
                                                          Similarity
                                                                                                                                                   biosynthesis;
                                                                                                       133
646 AA;
                                              Conservative
                                                                                                                               79
                                                                                                       22 G
83 G
136 G
; 73711 MW;
                       14
                                                         46.1%;
50.0%;
                                                                                                      Antibiotic resistance; GTP-binding; Plasmid.
22 GTP (BY SIMILARITY).
33 GTP (BY SIMILARITY).
66 GTP (BY SIMILARITY).
3711 MW; 1F3513BA2F644B6B CRC64;
                                                         Score 41; DB
Pred. No. 27;
                                              Mismatches
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sp_unclassified:*
sp_rvirus:*
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  0 Q9FXA2
Q9S4T6
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Q9U558
Q9N959
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Q928G4
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09cdg1 lactococcus
0928g4 listeria in
097sq3 streptococc
035745 mus musculu
0919t3 mus musculu
0912z3 mus musculu
091zz3 mus musculu
091zz3 mus musculu
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09n959 trypanosoma
09fxa2 arabidopsis
09s4t6 neisseria g
094iq3 anemia phyl
09m6e4 nicotiana t
057224 enterococcu
09ac90 clostridium
067709 gardnerella
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45	44	43	42	41	40	39	38	37	36	35	34	ω S	32	<u>3</u> 1	30	29	28	27	26	25	24	23	22	21	20	19	18	17
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Q62695 rattus norv	Q9quh9 mus musculu	Q9ue65 homo sapien	Q63419 rattus norv	P76578 escherichia	Q9vmb4 drosophila	Q9f4b2 bacillus st	Q9m549 cucumis sat		Q94gp3 oryza sativ	030707 streptomyce	Q9m6e5 nicotiana t	Q29136 tapirus ind	O14440 yarrowia li	Q9kuz7 vibrio chol	O23173 arabidopsis	Q93k56 fusobacteri	Q04134 drosophila		Q55136 synechocyst	5	Q9srk1 arabidopsis	P89471 herpes simp		P92948 arabidopsis	Q92qh2 rhizobium m		Q932i8 staphylococ	Q67710 gardnerella

ALIGNMENTS

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1	Query Match 53.9%; Score 48; DB 16; Length 709; Best Local Similarity 40.0%; pred. No. 11; Matches 6; Conservative 8; Mismatches 1; Indels 0; Gaps 0;	SEQUENCE /09 AA; 7/955 MW; LEAUZ9D65A/48032 CRC64;	proteome; GTP-binding.	PRINTS; PR00315; ELONGATNFCT.	Pfam; PF00679; EFG_C; 1.	InterPro; IPR000640; EFG_C. InterPro; IPR000795; GTP_EFTU.	P13551; 1DAR.		- I- SIMILARITY: BELONGS TO THE GTP-BINDING ELONGATION FACTOR FAMILY.	Genome Res. 11:731-753(2001).	lactis ssp. lactis IL1403.":	"The complete genome sequence of the lactic acid bacterium Lactococcus	Weissenbach J., Ehrlich S.D., Sorokin A.;	Bolotin A., Wincker P., Mauger S., Jaillon O., Malarme K.,	MEDLINE=21235186; PubMed=11337471;		SEQUENCE FROM N.A.	[1]	NCBI_TaxID-1360;		es; Bacillus/Clostridium group;	Lactococcus lactis (subsp. lactis) (Streptococcus lactis).	FUSA.	FACTOR G.	(TrEMBLrel. 19,	(TrEMBLrel. 17,	01-JUN-2001 (TrEMBLrel. 17, Created)		OSCOGI PRELIMINARY: PRT: 709 AA.	LT 1	

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SEQUENCE:
STRAIN-TIGR4;
STRAIN-TIGR4;
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MEDLINE-21357209; PubMed-11463916;
Tettelin H., Nelson K.E., Paulsen I.T., Eisen J.A., Read T.D.,
Tettelin H., Nelson K.E., Paulsen I.T., Haft D.H., Dodson R.J.,
Peterson S., Heidelberg J., DeBoy R.T., Haft D.H., Dodson R.J.,
Peterson S., Gwinn M., Kolonay J.F., Nelson W.C., Peterson J.D.,
Durkin A.S., Gwinn M., Kolonay J.F., Nelson W.C., Paterson J.D.,
Umayam L.A., White O., Salzberg S.L., Lewis M.R., Radune D.,
Holtzapple E., Khouri H., Wolf A.M., Utterback T.R., Hansen C.L.,
McDonald L.A., Feldblyum T.V., Angiuoli S., Dickinson T., Hickey
Holt I.E., Loftus B.J., Yang F., Smith H.O., Venter J.C.,
Holt I.E., Loftus B.J., Yang F., Smith H.O., Venter J.C.,
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Q97SQ3;
Q1-OCT-2001
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01-DEC-2001
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Bacillus/Staphylococcus
NCBI_TaxID=1642;
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01-DEC-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE 192 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=1313;
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N ELONGATION FACTOR G
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Pred. No. 3.5;
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Matches 9
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Spengler D., Villalba M., Hoffmann A., Pantaloni C.,
Spengler D., Villalba M., Hoffmann A., Pantaloni C.,
Bockaert J., Journot L.;
Submitted (JAN-1998) to the EMBL/GenBank/DDBJ databas
-! - SUBCELLULAR LOCATION: NUCLEAR (BY SIMILARITY).
EMBL; W95503; CAA64757.1; -.
EMBL; X95504; CAA64758.1; -.
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O35745;
O1-JAN-1998
O1-JAN-1998
O1-DEC-2001
                                                                                                                                                                                                                                                                                                                                                       SMART; SM00355; ZnF_C2H2; 7.

PROSITE; PS0028; ZINC_FINGER_C2H2_1; 7.

PROSITE; PS50157; ZINC_FINGER_C2H2_2; 6.

DNA-binding; Metal-binding; Nuclear protein; Zinc-finger.

SEQUENCE 667 AA; 75258 MW; 392F0AC979CBC10F CRC64;
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STRAIN-BALB/C; TISSUE-PITUITARY GLAND;
MEDLINE-97327559; PubMed-9184226;
Spengler D., Villalba M., Hoffmann A.,
Bockaert J., Journot L.;
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Pfam; PF00009; GTP_EFTU, 1.
Pfam; PF03144; GTP_EFTU_D2; 1.
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Mammalia; Eutheria;
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InterPro; IPR000795;
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EMBL; AE007340; AAK74451.1;
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6; Conserv
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Metazoa; Chordata; C
Metazoa; Rodentia; C
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(TrEMBLrel. 05, Last sequence update)
(TrEMBLrel. 19, Last annotation update)
PROTEIN REGULATOR OF APOPTOSIS AND CELL
                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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GTP_EFTU.
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                                                                                                                                                      . 0%;
                                                                                                      Score 45; DB Pred. No. 32; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the pituitary gland and the
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Pred. No.
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Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
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I'dentification of a splice variant of mouse Lotl gene.";

Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases.

R EMBL; AF324471; AAG48331.1; -.

R MGD; MGI:1100874; Zacl.

InterPro; IPR000822; Znf-C2H2.

R Pfam; pF00096; zf-C2H2; 7.

R SMART; SM00355; ZnF_C2H2; 7.

R SMART; SM00355; ZnF_C2H2; 7.

R PROSITE; PS00028; ZINC_FINGER_C2H2_1; 1.

R PROSITE; PS01078; ZINC_FINGER_C2H2_2; 6.

R PROSITE; PS0157; ZINC_FINGER_C2H2_2; 6.

R DNA-Dinding; Metal-Dinding; Zinc-finger.

Q SEQUENCE 675 AA; 76109 MW; 0F467D165CA5B097 CRC64;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q9JLQ4;
01-OCT-2000
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MEDLINE-20065118; PubMed-10597250;

Abdollahi A., Bao R., Hamilton T.C.;

"LOT1 is a growth suppressor gene down-regulated by t growth factor receptor ligands and encodes a nuclear
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
LOST ON TRANSFORMATION PROTEIN 1.
ZAC1 OR LOT1.
                                       SEQUENCE FROM N.A.

MEDILINE-20136063; PubMed-10669760;

Huang S.-M., Stallcup M.R.;

"Mouse Zacl, a Transcriptional Coactivator and Receptors.";

Mol. Cell. Biol. 20:1855-1867(2000).
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                       Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalla; Eutherla; Rodentia;
NCBI_TaxID=10090;
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nes 9; Conserv
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Last sequence update)
Last annotation updat
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 45;
Pred. No.
                                                                                                                                                                                                                                                                                                                                           Craniata; Vertebrata; Sciurognathi; Muridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                      Euteleostomi;
                                                                                                                                                                                                                                                                                                                                               Murinae;
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RESULT Q91ZZ3 ID Z92 ID Z93 ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Qγ
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Best Local
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Best Local :
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                                                                                                                                          Q9U558;
Q9U558;
01-MAY-2000
01-MAY-2000
01-DEC-2001
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01-DEC-2001
01-DEC-2001
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Q91ZZ3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MGD; MGI:1100874; Zacl.
InterPro; IPR00082; Znf-C2H2.
Pfam; PE0096; zf-C2H2; 7.
SMART; SM00355; ZnF_C2H2; 7.
SMART; SM00355; ZnF_C2H2; 7.
PROSITE; PS00028; ZINC_FINGER_C2H2_1; 7.
PROSITE; PS50157; ZINC_FINGER_C2H2_2; 6.
DNA-binding; Metal-binding; Zinc-finger.
DNA-binding; Metal-binding; Zinc-finger.
SEQUENCE 704 AA; 79214 MW; 3AF88262D504FA80 CRC64;
                                                                   Plasmodium vivax.
Eukaryota; Alveolata;
NCBI_TaxID=5855;
                                                                                                                                                                                                                                                                                                                                                                                                                     mouse beta-synuclein, a candidate for in neurodegeneration.";
Cytogenet. Cell Genet. 93:117-123(2001).
EMBL; AF348164; AAK83238.1; -
EMBL; AF348162; AAK83238.1; JOINED.
EMBL; AF348163; AAK83238.1; JOINED.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
             SEQUENCE FROM N.A. STRAIN-KPV97-73; Kim S.H., Kwon S.I
                                                                                                                 MEROZOITE SURFACE MSP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BETA-SYNUCLEIN.
                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sopher B.L., Koszdin K.L.,
Smith A.C., La Spada A.R.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
MEDLINE-21367613; PubMed-11474193;
"Analysis of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Genomic organization,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=10090;
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                                                                                                                                                                                                                                                                                        5 QPLLQVMMEPQGD 17 :||::::|||:||:
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9; Conserv
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                                                                                                                             0 (TrEMBLrel. 13, 0 (TrEMBLrel. 13, 1 (TrEMBLrel. 19, 1 (TrEMBLrel. 19, SURFACE PROTEIN 1
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                                                                                                                                                                                                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                     Conservative
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               S.I.;
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a candidate for involvement
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                                                                                                                             Created)
Last sequence update)
Last annotation updat
(FRAGMENT).
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Last sequence update)
Last annotation update)
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                 Score 44; I
Pred. No. 9
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                                                                                                                                                                                                     PRT;
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                                                                                     Haemosporida; Plasmodium
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34;
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Plasmodium vivax merozoite surface protein 1 gene

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RESULT
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OPEXA
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Best Loc
Matches
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Best Local
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01-OCT-2000 (TremBLrel. 15, C

01-OCT-2000 (TremBLrel. 15, L

01-OCT-2000 (TremBLrel. 15, L

POSSIBLE HYPOTHETICAL 14.6 KD

CHR1.04.
                              SEQUENCE FROM N.A.

Pederspiel N.A., Palm C.J., C
Altafi H., Araujo R., Huizar
Gonzalez A., Kremenetskaia I.
                                                                                                                                                                                                                                                                                                                             O9FXA2

O9FXA2;
O1-MAR-2001 (TrEMBLrel. 16, Created)
O1-MAR-2001 (TrEMBLrel. 16, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
PUTATIVE POLY-A BINDING PROTEIN.
F14J22.3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN-TREU927;
Hall N., Bowman S., Quail M., Ivens A.C., Kay Lennard N.J., Clark L.N., Harris B.R., Melvill Gerrard C., Rajandream M.A., Barrell B.G.;
Submitted (JUN-2000) to the EMBL/GenBank/DDBJ EMBL; AL359782; CAB95318];
SEQUENCE 227 AA; 26099 MW; AAF344E1959DDIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NON_TER
NON_TER
SEQUENCE
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Submitted
                                                                                                                                                                                                   Arabidopsis thaliana (Mouse-ear cress).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheopsermatophyta; Magnoliophyta; eudicotyledons; core eudicots; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Trypanosoma brucei.
Eukaryota; Euglenozoa;
NCBI_TaxID=5691;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL;
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Lee K.N., Song K
                                                                                                                                                                             NCBI_TaxID=3702;
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tted (OCT-1998) to the
AF098282; AAF25530.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity 57.88; Conservative
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9; Conser
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182 AA;
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; 19514 MW;
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56
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the EMBL/GenBank/DDBJ databases
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5, Last sequence upo
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6 KDA PROTEIN.
                           Conway A.B., Conn L., Hansen
r L., Rowley D., Buehler E., I
I., Kim C., Lenz C., Li J., Li
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., Melville S
B.G.;
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Liu S.,
                                       a N.F.,
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EMBL; AC011807; AAG13056.1; -.

EMBL; AF370517; AAK43894.1; -.

HSSP; P11940; 1CVJ.

Interpret: Tree-Cor.
                                                                        NON_TER
NON_TER
SEQUENCE
                                                                                                                                                                           Ferreira E., Louro D., Gomes J.P., Catry M.A., Vaz Pato M.V.; "Surveillance of tetracycline-resistant gonocci in Lisbon. of a TRNG strain with tetracycline MIC-2 mg/L."; Submitted (DEC-1998) to the EMBL/GenBank/DDBJ databases. EMBL; AF116348; AAD48368.1; -.
                                                                                                                                                                                                                                                                                                                                                                                      Q9S4T6;
                                                                                                                           Pfam;
                                                                                                                                                                                                                                                                              Bacteria; Proteobacteria; NCBI_TaxID=485;
                                                                                                                                                                                                                                                                                                                                 01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence up
01-DEC-2001 (TrEMBLrel. 19, Last annotation
TETRACYCLINE RESISTANCE PROTEIN (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pfam; PF00658; PABP; 1.

Pfam; PF00076; rrm; 4.

PRINTS; PR00961; HUDSXLRNA.

SMART; SM00517; PolyA; 1.

SMART; SM00360; RRM; 4.

SMART; SM00361; RRM, 1; 4.

PROSITE; PS01020; RRM; 4.

PROSITE; PS00030; RRM_RNP_1; UNKNOWN_1.

PROSITE; PS00030; RRM_RNP_1; UNKNOWN_1.
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Tones
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Luros S., Schwartz J., Shinn P., Toriumi M., Vysotskaia Walker M., Yu G., Ecker J., Theologis A., Davis R.W.; Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases
                                                                                                                Cyclin;
                                                                                                                                          Pfam;
                                                                                                                                                   InterPro; IPR000795; GTP_EFTU_D2.
                                                                                                                                                                                                                                           STRAIN-TRNG 400;
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PF03144; GTP_EFTU_D2; 1.
                                                                                                               GTP-binding.
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IPR002004; PABP.
IPR000504; RRM.
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29007 MW;
                        48.3%;
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46.78;
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Last sequence update)
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            Score 43; DB pred. No. 26; 3; Mismatches
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Pred. No. 48;
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                                                                                                                                                                                                                                                                                                subdivision;
                                                                         25BE46A2F2BDC067 CRC64;
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Matches 8
PROSITE; PS50102; RRM; 2.
PROSITE; PS00030; RRM_RNP_1; U.
NON_TER 1
SEQUENCE 470 ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q94IQ3;
01-DEC-2001
01-DEC-2001
01-DEC-2001
                                                                                            InterPro;
InterPro;
                                                                                                                 plants.";
Plant Sci. 152:101-114(2000).
EMBL; AF190657; AAF66825.1; -.
HSSP; P11940; ICVJ.
                                                                                                                                                                                                                 Nicotiana tabacum (Common tobacco).

Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Asteridae; euasterids I; Solanales; Solanaceae; Nicotiana.
                                                                                                                                                                                                                                                                       01-OCT 2000 (TrEMBLrel. 15, Created 01-OCT-2000 (TrEMBLrel. 15, Last se 01-DEC-2001 (TrEMBLrel. 19, Last an DOLY(A)-BINDING PROTEIN (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (APR-2001) to the EMBL/GenBank/DDBJ databases EMBL; AY030359; AAK51123.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vugrek O., Moepps B., Schraudolf H.;
"cDNA cloning of poly(A)-binding proteins
phyllitidis.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anemia phyllitidis (Fern)
Eukaryota; Viridiplantae;
Filicophyta; Filicopsida;
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                                               Pfam; PF00076; rrm; 2.
SMART; SM00517; PolyA; 1.
SMART; SM00360; RRM; 2.
                                                                                                                                                                       SEQUENCE FROM N.A.
Le H., Gallie D.R.;
                                                                                                                                                                                                                                                                                                                    Q9M6E4;
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                                                                                                                                                                                                      Asteridae; euasterids
NCBI_TaxID=4097;
                                                                                                                                                                                                                                                               PABP.
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                                                                                  Pfam; PF00658; PABP;
                                                                                                                                                               "Sequence
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les 8; Conserv
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IPR000504; RRM.
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        A2C3F46C3D688A12 CRC64;
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STRAIN=DS16; TRANSPOSON=TNYIU,
MEDLINE=92368175; PubMed=1323953;
Su Y.A., He P., Clewell D.B.;
Su Y.A., He P., Clewell D.B.;
"Characterization of the tet(M) determinant of Tn916: evidence regulation by transcription attenuation.";
arents Chemother. 36:769-778(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q57224;
Q57224;
01-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN-DS16: TRANSPOSON-TN916; MEDLINE-95207419; PubMed-789523; MEDLINE-95207419; PubMed-789523; Flannagan S.E., Zitzow L.A., Su Y.A., Clewell D.B.; "Nucleotide sequence of the 18-kb conjugative transposon Tn916 from Enterococcus faecalis.";
                                                                                                                                     Pfam; PF00009; GTP_EFTU; 1.
Pfam; PF03144; GTP_EFTU_D2; 1.
PRINTS; PR00315; ELONGATNECT.
                                                                                                                                                                                 InterPro; IPR000640;
InterPro; IPR000795;
InterPro; IPR004161;
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                                                                                                                                                                                                                                                                                                                                                                                                      Su y.A., Clewell D.B.;
"Characterization of the left 4 kb of "Characterization to the excision.";
determinants involved in excision.";
Plasmid 30:234-250(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE OF 636-639 FROM N.A. STRAIN=DS16; TRANSPOSON=TN916; MEDLINE=94134828; PubMed=8302931;
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Bacteria; Firmicutes;
Enterococcus.
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                                                                                                   GTP-binding
SEQUENCE (
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                                                                                                                                                                                                                                                    Submitted (MAY-1994) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                Clewell D.B.;
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                                                                                                                          PROSITE; PS00301;
                                                                                                                                                                                                                                                                           STRAIN-DS16; TRANSPOSON-TN916;
                                                                                                                                                                                                                                                                                       SEQUENCE FROM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=1351;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Plasmid
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IENPHPLLQTTVEP
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                                             Similarity 50.
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Bacillus/Clostridium group;
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                                            Score 43; DB
Pred. No. 67;
3; Mismatches
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(P SEQUENCE FROM N.A.

(C TRANSPOSON-CW459TET(M);

(X MEDLINE-21220755; PubMed-11320127;

(X MEDLINE-212207555; PubMed-11320127;

(X MEDLINE-212207555; PubMed-11320127;

(X ROBERTS A.P., Johanesen P.A., Lyras D., Mullany P., Rood J.I.;

(X Comparison of Th5397 from Clostridium difficile, Tn916 from

(Y Tenterococcus faecalis and the CW459tet(M) element from Clostridium

(Y Tenterococcus faecalis and the CW459tet(M) element from Clostridium

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(Y Tenterococcus faecalis and the CW459tet(M) element from Clostridiu
                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 50.0
Matches 7; Conservative
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09AC90; O1-JUN-2001 (TrEMBLrel 17, Created)
01-JUN-2001 (TrEMBLrel 17, Last sequence update)
01-DEC-2001 (TrEMBLrel 19, Last annotation update)
TET(M).
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Bacteria; Firmicutes; Bacillus/Clostridium group; Clostridiaceae; Clostridium.
                                                                                                                                                                                                                                              1 LQTPQPLLQVMMEP 14 :: | | | | | : | | 352
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